SEQUENCE LISTING

<110> OOE, Norihasa MATSUNAGA, Haruyuki <120> CELL FOR MEASURING THE ABILITY TO CONTROL THE ACTIVITY OF A LIGAND-RESPONSIVE TRANSCRIPTION CONTROL FACTOR <130> 2185-0424P <140> 09/550,173 <141> 2000-04-14 <150> JP H11-106791 <151> 1999-04-14 <150> JP H11-106792 <151> 1999-04-14 <150> JP H11-106793 <151> 1999-04-14 <150> JP H11-107774 <151> 1999-04-15 <160> 34 <170> PatentIn Ver. 2.1 <210> 1 <211> 6 <212> DNA <213> Unknown Organism <220> <223> Description of Unknown Organism: consensus sequence of a dioxin-responsive sequence <220> <221> Unsure <222> (1)..(1) <223> n = t or a <300> <303> J. Biol. Chem. <304> 271 <306> 3952-3958 <307> 1996-02-01 <400> 1 ngcgtg <210> 2 <211> 16 <212> DNA

<213> Unknown Organism

6

<220> <223>	Description of Unknown Organism: consensus sequence of an estrogen-responsive sequence	
<222>	Unsure (7)(9) n = a,c,g,t any unknown or other.	
<400> aggtc	2 annnt gacctt	16
<210><211><211><212><213>	20	
<220> <223>	Description of Artificial Sequence:primer for PCR with human genomic DNA.	
<400> ttgag	3 ctagg cacgcaaata	20
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<220> <223>	Description of Artificial Sequence:primer for PCR with human genomic DNA	
<400> gcttt	4 gattg gcagagcaca	20
<210><211><212><212><213>	51	
<220> <223>	The sequence is composed of nucleotide sequences derived from a nucleotide sequence near the TATA box of a mouse metallothionein I gene. The sequence is introduced into mouse and human cells.	
<400> gatct	5 cgact ataaagaggg caggctgtcc tcaagcgtca ccacgacttc a	51
<210> <211>		

	<212> <213>	mouse	
	<220> <223>	The sequence is composed of nucleotide sequences derived from a nucleotide sequence near the TATA box of a mouse metallothionein I gene. The sequence is introduced into mouse and human cells.	
	<400> agctt	6 gaagt cgtggtgacg cttagaggac agcctgccct ctttatagtc ga	52
	<210><211><211><212><213>	33	
	<220> <223>	The sequence is located at the upstream of a Xenopus-derived vitellogenin gene containing a recognition sequence of an estrogen receptor. The sequence is introduced into mouse and human cells.	
	<400> tcgaca	7 Baagt caggtcacag tgacctgatc aag	33
	<210><211><211><212><213>	31	•
	<220> <223>	Description of Artificial Sequence:primer for PCR with pTK beta	
	<400> cggcag	8 gatet tetttagtte tatgatgaca e	31
	<210><211><211><212><213>	29	
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	<400> cggaag	9 Jottg atotgoggoa ogotgttga	29
•	<210><211><211><212>	35	

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<220; <223;	> > Description of Artificial Sequence:primer for PCR with human cDNA	
<400 cctg	> 10 egggga caeggtetge accetgeeeg eggee	35
<2203 <2233	Description of Artificial Sequence:primer for PCR with human cDNA	
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<210><211><211><212><213>	40	
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<400> caggg	13 agctc tcagactgtg gcagggaaac cctct	35
<210><211><211><212><213>	35	
<220> <223>	Description of Artificial Sequence:primer for PCR with human cDNA	

<400> 14 ttgagttact gagtccgatg aatgtgcttg ctctg	35
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<400> 16 gccgcggccg cccagccacc atggatataa aaaactcacc atctagcctt aattc	55
<pre> <210> 17 <211> 43 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence:primer for</pre>	
adding Kozak consensus sequence to human cDNA <400> 17 gggtctagaa atgagggacc acacagcaga aagatgaagc cca	43
<210> 18 <211> 52 <212> DNA <213> mouse	
<pre><220> <223> The sequence is derived from a nucleotide sequence near the TATA box of a mouse metallothionein I gene. The sequence is used for human cells. The sequence is introduced into human cells.</pre>	
<400> 18 gatctcgact ataaagaggg caggctgtcc tctaagcgtc accacgactt ca	52

<220> <223>	The sequence is derived from a nucleotide sequence near the TATA box of a mouse metallothionein I gene. The sequence is used for human cells. The sequence is introduced into human cells.	
<400> agctt	· 19 gaagt cgtggtgacg cttagaggac agcctgccct ctttatagtc ga	52
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<211> 35 <212> DNA <213> Artificial Sequence	
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adding Kozak consensus sequence to human cDNA	
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<210> 24 <211> 38 <212> DNA <213> mammal	
<220> <223> The sequence is containing a recognition sequence (TRE) of a thyroid receptor. The sequence is introduced into human cells.	
<400> 24 caaggggatc cagcttgacc tgacgtcagg tcaagtcg	38
<210> 25 <211> 52 <212> DNA <213> mouse	
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	with human cDNA	
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tggcc	geetg aggetttaga etteetgate eteaa	35
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<400>	• 31			
ttact	aacct ataaccccca acagtatgac agaaa	35		
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<400>	33			
cccago	ccacc atgacagaaa atggccttac agcttgggac	40		
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<400>	34			
cagtct	aatc ctcgaacact tccaggaaca aaggg	35	•	

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